CRF Problem Report

The Scientific and Technical Information Center (STIC) experienced a problem when processing the following computer readable form (CRF):

Application Serial Number: ___09/8336735___

Filing Date: ___~~10/90/01~~___ 04-13-2001

Date Processed by STIC: ___10/90/01___

STIC Contact: **Mark Spencer, 703-308-4212**

Nature of Problem:

The CRF (was):

____ (circle one) Damaged or Unreadable (for Unreadable, see attached)

____ Blank (no files on CRF) (see attached)

____ Empty file (filename present, but no bytes in file) (see attached)

____ Virus-infected. Virus name: _____ The STIC will not process the CRF.

____ Not saved in ASCII text

____ Sequence Listing was embedded in the file. According to Sequence Rules, submitted file should **only** be the Sequence Listing.

__✓__ Did not contain a Sequence Listing. (see attached sample)

____ Other:

PLEASE USE THE CHECKER VERSION 3.0 PROGRAM TO REDUCE ERRORS. SEE BELOW FOR DETAILS:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

http://www.uspto.gov/web/offices/pac/checker

RAW SEQUENCE LISTING   DATE: 10/30/2001
PATENT APPLICATION: US/09/833,675   TIME: 12:32:22

          Input Set : A:\PTO.mh.txt
          Output Set: N:\CRF3\10302001\I833675.raw

    3 <110> APPLICANT: Aventis Behring GmbH
    5 <120> TITLE OF INVENTION: Process for finding oligonucleotide sequences
    6   for nucleic acid amplification methods
    8 <130> FILE REFERENCE: 2000/A006-A3b
C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/833,675
C--> 11 <141> CURRENT FILING DATE: 2001-04-13
E--> 13 <160> NUMBER OF SEQ ID NOS: 9
    15 <170> SOFTWARE: PatentIn Ver. 2.1

ERRORED SEQUENCES

Does Not Comply
Corrected Diskette Needed

See Page 2 of 3 A

Deleted Sequence
Between Sequence
Non AsCII text

<210> 1
<211> 27
<212> DNA
<213> Red sea bream iridovirus gene

<210> 2
<211> 18
<212> DNA
<213> Homo sapiens

Actual file contents
As of 10/30/01
09/833675   mot

Sequence Listing Includes
no fields 400 nor any gene
sequences

Therefore field 160 Input 9
Found 0 — Errored

<210> 3
<211> 22
<212> DNA
<213> Homo sapiens

<210> 4
<211> 21
<212> DNA
<213> Human DNA sequence from PAC 388M5

There are no genetic sequences on the
sequence listing.

**The type of errors shown exist throughout
the Sequence Listing. Please check subsequent
sequences for similar errors.**

VERIFICATION SUMMARY   DATE: 10/30/2001
PATENT APPLICATION: **US/09/833,675**   TIME: 12:32:23

Input Set : **A:\PTO.mh.txt**
Output Set: **N:\CRF3\10302001\I833675.raw**

```
L:10 M:270 C: Current Application Number differs, Replaced Application Number
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:24 M:280 W: Numeric Identifier already exists, Length not replaced.
L:25 M:280 W: Numeric Identifier already exists, Type not replaced.
L:26 M:280 W: Numeric Identifier already exists, Organism not replaced.
L:34 M:280 W: Numeric Identifier already exists, Length not replaced.
L:35 M:280 W: Numeric Identifier already exists, Type not replaced.
L:36 M:280 W: Numeric Identifier already exists, Organism not replaced.
L:40 M:280 W: Numeric Identifier already exists, Length not replaced.
L:41 M:280 W: Numeric Identifier already exists, Type not replaced.
L:42 M:280 W: Numeric Identifier already exists, Organism not replaced.
L:47 M:280 W: Numeric Identifier already exists, Length not replaced.
L:48 M:280 W: Numeric Identifier already exists, Type not replaced.
L:49 M:280 W: Numeric Identifier already exists, Organism not replaced.
L:54 M:280 W: Numeric Identifier already exists, Length not replaced.
L:55 M:280 W: Numeric Identifier already exists, Type not replaced.
L:56 M:280 W: Numeric Identifier already exists, Organism not replaced.
L:63 M:280 W: Numeric Identifier already exists, Length not replaced.
L:64 M:280 W: Numeric Identifier already exists, Type not replaced.
L:65 M:280 W: Numeric Identifier already exists, Organism not replaced.
L:70 M:280 W: Numeric Identifier already exists, Length not replaced.
L:71 M:280 W: Numeric Identifier already exists, Type not replaced.
L:72 M:280 W: Numeric Identifier already exists, Organism not replaced.
L:77 M:280 W: Numeric Identifier already exists, Length not replaced.
L:78 M:280 W: Numeric Identifier already exists, Type not replaced.
L:79 M:280 W: Numeric Identifier already exists, Organism not replaced.
L:13 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (9) Counted (0)
```